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Copyright (c) 1993 - 2000 Compugen Ltd
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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                                 cc a 53 kba BRCA1 modulator protein (see AAW37881) that binds to the ct tumour suppressor gene product BRCA1, and which is characterised by cc a zinc finger domain and a leucine zipper motif. 3 cDNA clones cc (see also ANV29063 and ANV29064) coding for BRCA1 modulator proteins cc yeast two-hybrid assay with a GAL4-BRCA1(8-1293) fusion as bait. cc vectors and host cells comprising the isolated nucleic acid sequences are claimed, as well as a process for producing BRCA1 cc modulator protein by culturing these host cells. BRCA1 modulator cc proteins and nucleic acids can be used to diagnose diseases compounds that alter BRCA1 interaction with BRCA1 modulators for the treatment of such diseases.
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                                    AAS27719
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                  AAS27719;
                                                                              GGACCAGAGTGCTTGAGGCATCTCAGGCAGCCTCAGCCCCAAGCTTCTACCTGCCTTTGAC
                                                                                                                                                                                        AGACTGCTCTTCCTGCCCTCACCCCTGCCCCACTCCTACGACTGGGGAGCTGACCATGACCAG
                                                                                                                                                                                                                                                                                                  TTCGGGACCAGCCTGAGGTGTAAGGGCAGACAAACAGGTGAGGGTGAGTGTGACACCCAG
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                                                                                                                           TTGCTTCTA-GCATAGCCTGGGCCAAGCAGGGTGGGGGAATGGAGATAGACATGGGATGT
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                                                                                                                  ttgcttctaggcatagcctgggccaagcagggtggggaatggaggatag-catgggatgt
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                                    standard; DNA; 148
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DNA
encoding
 novel
signal transduction
pathway
protein,
Seq
ID
 1379.
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Neuroprotective; cytostatic; dermatological; immunosuppressive; tumous antiinflammatory; anti-HIV; antibacterial; antiinflammatory; cancer; immune system disorder; rheumatoid arthritis; inflammatory condition; organ transplant rejection; infection; hepatitis C; blood disorder; sickle cell anaemia; hyperproliferative disorder; Gaucher's disease; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; chromosomal abnormality; Down syndrome; ischaemia; renal disorder; cardiovascular; respiratory; wound healing; endocrine; Addison's disease; productive system; gastrointestinal; liver disorder; AIDS; ds; acquired immune deficiency syndrome. Addison's disease; AIDS; ds; tumour;

Homo

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31 JAN - 2000
04 FEB - 2000
24 FEB - 2000
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16 - MAR - 2000
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19 - MAY - 2000
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                                     2000US-0180628
2000US-0180528
2000US-0180564
2000US-019076
2000US-0199123
2000US-0205467
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2000US-0232397.
2000US-02323400.
2000US-0233400.
2000US-0233063.
2000US-0233064.
2000US-0233064.
2000US-0234025.
2000US-0234025.
2000US-0234997.
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2000US-0234988.
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2000US-023498.
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2000US-0236370. 2000US-0236802. 2000US-0237037. 2000US-0237038. 2000US-0237039.

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RESULT
AAA30290
ID AAA3
XX
AC AAA3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to novel isolated polypeptides (I), and comply provides (II). (II) and the antibody to (I) are useful for CC disorders (e.g. congenital and acquired immunodeficiencies, autoimmune CC disorders (e.g. congenital and acquired immunodeficiencies, autoimmune CC transplant rejections and graft versus host disease, infectious diseases CC (e.g. hepatitis C), bleeding disorders, haemoglobin abnormalities and CC other blood-related disorders (sickle cell anaemia), myeloproliferative CC disorders (e.g. Gaucher's disease and cancer), neurodegenerative CC disorders (e.g. Gaucher's disease, Parkinson's disease), chromosomal CC disorders (e.g. glomerulonephritis), cardiovascular disorders (e.g. glomerulonephritis), cardiovascular disorders (e.g. glomerulonephritis), cardiovascular disorders (e.g. dalcheila) cell proliferation, endocrine disorders (e.g. addison's disease), reproductive system disorders, gastrointestinal CC disorder (inflammatory disorders, liver disorders, gastrointestinal CC disorder (inflammatory disorders), liver disorders, cirrhosts), CC as stimulators of B-cell responsiveness to pathogens, activators of CC T-cells, to induce higher affinity antibodies, and as a means to induce CC syndrome (AIDS). AAS26976-AAS27850 represent novel signal transduction CC pathway protein coding sequences and PCR primers of the invention.
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Best Local Similarity 100.0%;
Matches 148; Conservative
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                                AAA30290;
                                                                                      AAA30290 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; SEQ ID No 1379; 880pp; English.
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2000US-025719.
2000US-0251479.
2000US-0251856.
2000US-0251868.
2000US-0251869.
2000US-0251990.
2000US-0251990.
2000US-0254097.
2001US-0259678.
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2000US-0251030
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Pred. No. 3e-30;
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26-SEP-2000
27-SEP-2000
29-SEP-2000
20-CCT-2000
21-NOV-2000
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2000US-0237040.
2000US-0239935.
2000US-0239935.
2000US-0241786.
2000US-0241786.
2000US-0241809.
2000US-02446174.
2000US-0246475.
2000US-0246476.
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2000US-024661.
2000US-0249218.
2000US-0249219.
2000US-0249218.

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Query Match
Best Local S
Matches 139
                                                                                                efficient persistence of rhadino virus DNA in mammalian cells. Persistent rhadino virus infection is implicated in a variety of diseases e.g. Kaposi's Sarcoma (KS), Primary Effusion Lymphoma (PEL) and multicentric Castleman's disease. In addition, KS is a common malignancy in HIV patients. KSHV persists in host cells in a latent form. One of the few genes expressed from the latent viral DNA is LANA. LANA associates with both human chromosomes and with the rhadino virus cis-acting element (RVCAE), thereby providing a tethering function: the KSHV DNA episome is "tied" to the host chromosomes. This allows the viral DNA to persist in the host cell. The present sequence may be used to screen and identify molecules that inhibit LANA interaction with RVCAE, thereby interfering with the latency cycle of this virus. Potential antiviral treatments for the above mentioned diseases may therefore be based on LANA deregulation.
                                                                                                                                                                                                                                                                                          The present sequence is the Kaposi's sarcoma-associated herpesvirus, (KSHV) latency-associated nuclear antigen (LANA) gene. KSHV is also as Human Herpes Virus 8 (HHV8) and belongs to the rhadino virus, or gamma-2 herpes virus class. The LANA protein is necessary for the
                                                                        Sequence 3489
                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 6;
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                                                                                                                                                                                                                                                                                                                                                                                                            Treating or preventing a disease infection in a mammal which inclu
                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (KIEF/) KIEFF E D. (BALL/) BALLESTAS (KAYE/) KAYE K M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-NOV-1998;
21-APR-1999;
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    139;
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                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY96255
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   Conservative
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/note= "nuclear localisation signal,
190..210
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                2.6%;
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                                                                                                                                                                                                                                                                                                                                                                  English.
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; Score 52; DB;
; Pred. No. 0.00;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                           sease associated with rhodino virus includes Kaposi's Sarcoma and Prim
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                                                                        1137 G;
                                                                        437
               DB 21;
   145;
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                            Length 3489;
 Indels
                                                                        other;
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 0;
Gaps
 0;
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505 CTGAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAAGAG 564

the Kaposi's

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RESULT
AAF82901
                   tethering protein is LANA. The composition is useful in aiding the retention of the viral DNA in the host cell. The viral vector encodes a protein suitable for tethering DNA to Histone H1. Methods for screening for compounds which are agonistic or antagonistic for the tethering of viral proteins to histone H1 and DNA binding sites are useful for developing the method of viral transfer. The composition has applications to gene therapy, including the treatment of multiple sclerosis, Parkinson's disease, Huntington disease and diabetes. The present
                                                                                                                                                                                                      The invention provides a composition comprising nucleic acid, histone H. protein and expression vector operationally encoding a protein suitable for tethering the nucleic acid to the histone HI protein, where the tethering protein is LANA. The composition is useful in aiding the
                                                                                                                                                                                                                                                                                                                                                                                                                       A composition for use in that includes a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
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sequence represents the nucleotide
                                                                                                                                                                                                                                                                                                                                              Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                    protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Robertson ES,
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kaposi's sarcoma; acquired immune deficiency syndrome; AIDS; DHFR; Bcl-idihydrofolate reductase; LUR; long unique region; vaccine; prophylaxis; diagnosis; treatment; HHV8; capsid protein IV; tegument protein IV; glycoprotein; kaposin; cyclin D; immediate early protein; IEP; OX-2; v-adh; G-protein coupled receptor; FGARAT; ds.
This sequence is a fragment of the Kaposi's sarcoma-associated herpesvirus (KSHV) LUR (long unique region). This fragment contains coding regions for ORF65 which encodes capsid protein IV, ORF66, ORI which encodes tegument protein IV, ORF68 which encodes a glycoprote:
                                                                                                                          Kaposi's sarcoma-associated herpes virus di:hydro:folate reductase and is useful or diagnosis of Kaposi's sarcoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV73805 standard; DNA; 32207
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                                                                                          Disclosure;
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(LANA), which acts as
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48.9%;
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Pred. No.
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for treatment, p
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                                                                                                                                                                                                                                                                                                                        status;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP;
              /product= macrophage inflammatory
complement (27137..27424)
                                                                  /product= interleukin 6
complement (21548..21832)
                                                                                                                        complement (17261..17875)
                                                                                                                                                                             /product= complement-binding
8699..11236
                                                                                                                                          /product= glycoprotein
                                                     /*tag=
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19785 CAGGATGAGCAGCAGCAGCAGGATGAGCAGCAGCAGGAGGAGCAGCAGGAG
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                                                                                                                                                                                                               KSHV; HHV8; human herpes virus 8; macrophage inflammatory protein II; interleukin-6; IL-6; interferon regulatory factor; rheumatoid arthritis, complement-binding protein; glycoprotein; capsid protein IV; infection; immediate early protein; Kaposi's sarcoma; protective vaccine; lymphoma.
                                                                                                         Kaposi's sarcoma-associated herpes virus.
                                                                                                                                                                                     lymphoproliferative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         505 CTGAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAAGAG
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/*tag=
                                                     Location/Qualifiers
                                                                                                                                                            anti-inflammatory agent;
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Pred. No. 0.00
0; Mismatches
                                                                                                                                                            leukaemia; splenomegaly; mycosis
ammatory agent; therapy; ds.
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This sequence represents the long unique region and terminal repeat of the Kaposi's sarcoma-associated herpes virus (KSHV). KSHV is also known as human herpes virus 8 (HHV8). This sequence contains the DNAs of the invention which encode KSHV polypeptides selected from: (a) viral C macrophage inflammatory protein (MIP) II; (b) viral interleukin-6 (IL-6); (c) viral IRF 1; (d) complement-binding protein; glycoproteins B, M or L; (d) capsid protein IV encoded by ORF65; and (e) immediate early protein encoded by ORF65; and (e) immediate early protein sencoded by it, and antibodies (Ab) specific for the nucleic acid, proteins encoded by it, and antibodies (Ab) specific for the proteins are useful for detecting HHVB, specifically for diagnosis of Kaposi's sarcoma, in body fluids or tissue samples. HHVB infections can be treated with antisense or triplex forming molecules or agents that bind specifically to the protein. Ab may be used for prophylaxis or treatment of HHVB infection, while the protein can be used in protective vaccines. Ab may also be used
                                                                                                                                                                                                                                                                                                                                                                     25-JUL-1996;
25-JUL-1996;
25-JUL-1996;
05-SEP-1996;
10-OCT-1996;
13-NOV-1996;
13-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-NOV-1996;
25-JUL-1996;
25-JUL-1996;
                                                                                                                                                                                                                                     New nucleic acid encoding Kaposi's sarcoma associated herpes proteins - useful for, e.g. detecting levels of HHV8 in, and preparation of vaccines for treatment of, HIV patients
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96US-0687253.
96US-0688814.
96US-0708678.
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96US-0747887.
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89600..90541
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complement (111931..112443)
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complement (93636..94127)
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90173..90643
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lymphomas,
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t (123808..127296)
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t (88410..88910)
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t (58976..60175)
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                                                                                                                                                                                                                                                                                                                    PS,
                                                                                                                                                                                                                                   of HHV8 in,
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be implicated
             also be used
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g. treating frames.
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                                                                                                                                                                          13-APR-2001;
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                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                lymphocytic leukaemia; lymphoma; plasmocytoma; Hodgkin's disease; glioma; ss.
                                                                                                                                                                                                                                                                                                                                                     Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 137507 BP;
                                                                                                                     (MILL-) MILLENNIUM PREDICTIVE
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Novel using

nucleic acid, used as TAXOL to treat cancer

cell grow

erker to

determine the effectiveness in individuals -

Claim 1; Page 450-451; 527pp; English

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         developing treatments for cancer, particularly those cancers which display resistance to agents and exhibit expression of the markers. The anticancer agents developed by the novel method can be used to treat cancer. Probes based on the markers can be used to detect transcripts or genomic sequences corresponding to the markers, in the identification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cells are shown not to express one of the 804 resistance markers. The methods can be used to determine the effectiveness of TAXOL in the treatment of cancer cell growth in an individual. The markers can be used as targets in developing anti-cancer agents such as chemotherapeutic compounds. The markers can also be used as targets
                                complement related protein; cytochrome; kinesin; cytokinterleukin; G-protein coupled receptor; thioesterase; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        1005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cells or tissues which mis-express the protein. Cancers which may be targeted include carcinoma (e.g. squamous cell carcinoma), sarcoma (e.g. fibrosarcoma) leukaemia (e.g. lymphocytic leukaemia) lymphoma, plasmocytoma, reticulum cell sarcoma, Hodgkin's disease tumours (e.g. glioma). The present sequence is one of the 1046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to 1046 novel nucleic acids which are used as markers for determining the sensitivity of a cancer cell to the anticancer agent TAXOL. Cancer cells can be treated with TAXOL when they are shown to express one of the 242 sensitivity markers or the
                                                                                   neuroprotective; antimicrobial; gene therapy; vaccine; amylase; can amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating f
                                                                                                                                                                                                                                                                                         AAL30971
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                                                                                                                                                                                                                   24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     novel cancer cell markers.
                                                                                                                                          Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -  \frac{1}{2} \left( \frac{1}{2} \right) = \frac{1}{2} \left( \frac{1}{2} \right) \left( \frac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                 Human; nootropic; immunosuppressant; cytostatic; gene therapy; cance peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1885
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shimkets RA,
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                                                                                                                  Homo sapiens.
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19-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7453
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CATGGA 602
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DB; AAM39213.
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                                     gctggacgacctcacggtggacctggaccaccagcgccaggtcgcctccaacttggagaa
                                                                         GCAGGATGAGACCAAACAAGCACAAGAGGAGGCGGGCCGGCTCAGGAGCAAGATGAAGAC 596
                                                                                                               ggaggagaaggcactggcgtatgacaaactggagaagaccaagaaccgcctgcagcagga
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                                                                                                                                                                                           aagtctggaagaagccaagaagcttctgaaggacgcggaggccctgagccagcgcct
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Query Match Best Local Similarity

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25-APR-2000;
09-JUL-2000;
19-JUL-2000;
                                immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, Activiny, inhibin activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tang
Wang
                                                                                                                                                                                                                                                                                                                Claim 1;
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                        .N.S disorders
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DB; AAM39214.
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Wang Z,
                                                                                                                                                                                                                                                                                                              SEQ ID NO 573; 10078pp; English
sequence
                                                                                                                                                                                                                                                      on relates to human nucleic acids (AAI57798-AAI61369) polypeptides (AAM38642-AAM42213) with nootropic,
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Wehrman T,
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        nucleic acids and polypeptides, useful
as central nervous system injuries -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide in gene therapy be used to treat diseases of the peripheral nervous of the invention may be used to treat diseases of the peripheral nervous of the peripher
       4315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               specification.
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ote: The sequence data
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                                                                            CACTCTGCGGGATACGCTGGAAGAACGCAATGCTACTGTGGTATCTCTGCAGCAGGCCTT
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aagtctggaagaagccaagaagcttctgaaggacgcggaggccctgagccagcgcct
                                                                                                                                                 cctgcagtcccagttggctgataccaagaagaaagtagatgacgacctgggaacaattga
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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RESULT 1
                                                                                                    Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; gene therapy; cance amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening.
26-JUL-2001
                                   WO200153312-A1
                                                                                                                                                                                                                              Human polynucleotide SEQ ID NO 4145.
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Best Local Similarity
Matches 145; Conserv
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25-APR-2000;
09-JUL-2000;
19-JUL-2000;
19-JUL-2000;
14-SEP-2000;
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                                                                                                                                                                                                                                                                                    4195 tcaggagcagcaggaggaggaggaggaggccaggaagaacctggagaagcaagtgctggc
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                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 7741 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acids and polypeptides, useful such as central nervous system injuries -
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29-NOV-2000;
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                                                                                                                                                                                                    417
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gaagca
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                            CATGGA 602
                                                                                                                              GGGCAAGGCCGAGATGCTGTGCTCCACACTGAAAAAGCAGATGAAGTACTTAGAGCAGCA
                                                                                                                                                                                      CACTCTGCGGGATACGCTGGAAGAACGCAATGCTACTGTGGTATCTCTGCAGCAGGCCTT
                                                                                                                                                                                                                             cctgcagtcccagttggctgataccaagaagaagtagatgacgacctgggaacaattga
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                                                      gctggacgacctcacggtggacctggaccaccagcgccaggtcgcctccaacttggagaa 4494
                                                                                 ggaggagaaggcactggcgtatgacaaactggagaagaccaagaaccgcctgcagcagga
                                                                                                                                                                    aagtctggaagaagccaagaagcttctgaaggacgcgggaggccctgagccagcgcct
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Wang Z,
Zhou P,
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2000US-0552317.

2000US-0598042.

2000US-0620312.

2000US-0653450.

2000US-0662191.

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2000US-0727344.
                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                  2303 A; 1751 C; 2080 G; 1606 T;
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Goodrich R,
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Wehrman T,
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Xu C, Xue
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Xu C, Xue AJ,
Drmanac RT;
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                                                                                                                                                                                                                                                                                                                                                                      DB 22;
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Yang Y,
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482 AGGCCGAGATGCTGTGCTCCACACTGAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGG 541

agctggagaggagcttgtccgaactcaaaaaccagatgggtaagatggggctgctacaag

Matches

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al Similarity 49.1 152; Conservative

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DB 23;).027;

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Sequence

5154

BP;

1544 A; 1236

c; 1478 G;

896 Ŧ; 0

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RESULT 14
AAS84859
                  CC polynucleotides are also used in diagnostics as expressed sequence tags CC for identifying expressed genes. (I) is useful in gene therapy techniques CC to restore mormal activity of (II) or to treat disease states involving CC (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as CC a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating CC disorders involving aberrant protein expression or biological activity. CC The polypeptide and polynucleotide sequences have applications in CC diagnostics, forensics, gene mapping, identification of mutations CC responsible for genetic disorders or other traits to assess biodiversity amino acid sequences. AAS64197-AAS94564 represent novel human CC diagnostic coding sequences of the invention.

CC Note: The sequence data for this patent did not appear in the printed constitution, but was obtained in electronic format directly from WIPO as ferromences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromc and gene mapping, and in recombinant production of (II). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated polynucleotide and encoded polypeptides, useful diagnostics, forensics, gene mapping, identification of mutauresponsible for genetic disorders or other traits and to asserted.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1;
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23-AUG-2000; 2000US-0649167
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ftp.wipo.int/pub/published_pct_sequences
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         \begin{array}{c} \texttt{CCCCCCCCX} \\ \texttt{X} \\ \texttt{Z} \\ 
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making
                                                                        A process has been developed for the production of a DNA fragment encoding silk protein. The process involves: (a) selecting target DNA, from a silk-producing spider, that contains many repetitive and non-repetitive regions; (b) selecting a single-stranded DNA primer of at least 10 nucleotides with a sequence that is complementary to a region of the target; (c) repetitively combining the primer with melted target DNA, incubating the mixture with nucleotides and a DNA polymerase with
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proofreading activity to produce a DNA fragment which is comp
to the target and is at least 2 kb long. The present sequence
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26-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               represents a 2 Kb DNA sequence which encodes the spider silk protein from Mephila clavipes. The DNA fragment can be used to make fibres, films, woven articles, e.g. for use in parachutes, sails, body armour, and absorbers (e.g. of heavy metals, biological weapons, DNA, chemicals flavours and fragrances). The high molecular weight (90-250 kD) of spider silk proteins can be produced on a commercial scale (at over 2 g/1 cell mass). It has better tensile strength and elasticity than silkworm silk. Inclusion of both repetitive and non-repetitive regions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cagccgctgcagcagctggtggtgccggacaaggaggatatggaggtcttggaaaccaag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAAGAGGAGG-CGGGCCGGCTCAG
                                                                                                                                                                                                       foetal liver single exon nucleic acid probe
                                                                                                                                                                            foetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2004 BP; 481 A;
2000US-0180312.
2000US-0207456.
2000US-0608408.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                          2001WO-US00669
                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.4%;
                                                                                                                                                                           gene
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                                                                                                                                                                            expression;
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Pred.
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No. 0
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0.025;
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DNA, chemicals,
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RESULT 17
ABA27737/C
ID ABA277
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ABA277
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ABA277
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DT 23-JAN
XX
Human,
KW Human,
KW Cardio
KW Congen
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Congen
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Matches 100
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21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
 04-FEB-2000; 2000US-0180312
                                                                                                                                                     Human; gene expression; heart; microarray; vascular system;
cardiovascular disease; hypertension; cardiac arrhythmia;
                                                                                                                                                                                                                                                                            737/c
ABA27737 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Note: The sequence data for this patent did not form part of printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a single exon nucleic acid probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting,
                           30-JAN-2001; 2001WO-US00666
                                                       09-AUG-2001
                                                                                   WO200157274-A2
                                                                                                                                        congenital
                                                                                                                                                                                                                         23-JAN-2002
                                                                                                                                                                                                                                                   ABA27737;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 7124; 639pp + sequence listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    analyzing
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                                                                                                                                                                                                                                                                                                                                                                             GTGGAACAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          genome-derived single exon nucleic acid |
zing gene expression in human fetal liver
                                                                                                                                                                                              #6203
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                                                                                                                                         heart disease;
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ilarity 52.9%;
Conservative
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2000US-0234687.
2000US-0236359.
2000US-0024263.
                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                            690
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                                                                                                                                                                                                                                                                              DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , 68
                                                                                                                                                                                             heart cell sample
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30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                               measuring human gene expression in a sample derived from human heart. present sequence is one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derive from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular systems. Cardiovascular disease, hypertension, cardiac arrhythmias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Single
hearts
                                                                                                                                                                                                                                                                                                                                                                                                    Note: The sequence data for this patent did not form part of the specification, but was obtained in electronic format directly from
                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                      at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                       congenital heart disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-OCT-2000;
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 184
                               682 GTGGAACAG 690
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                                                                                                622 CAGAGCCAGCTCCCTGAGGTGGAGGAGATGATCCGAGACATGGGTGTGGGACAGTCAGCG
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                                                                                                                                                                  562 GAGGAGGCGGCCGGCTCAGGAGCAAGATGAAGACCATGGAGCAGATTGAGCTTCTACTC 621
                                                                                                                                                                                                   502 ACACTGAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAA 561
GAGGAGGAG
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                                                                                                                                                                                                                                                                                                                                                    475 BP;
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                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention relates to single exon nucleic acid probes
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2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-02346359.
2000US-0236359.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6203;
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Pred. No. 0.01:
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                      194 T; 0 other;
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ARX66973/C
ID AAK069
XX AAK069
XX AAK069
XX O5-NOV
DT 05-NOV
DE Human
XX microa
KW Human;
KW microa
KW epilep
XX epilep
XX PN WO2001
XX
                                                                 Human; brain expressed exon; gene expressi microarray; Alzheimer's disease; multiple
                                 Homo sapiens
                                                                                                                           05-NOV-2001
                                                                                                                                                 AAK06973;
                                                                                                                                                                    AAK06973 standard;
                                                                                                   brain expressed single
                                                       cancer;
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                                                       SS
                                                                                                                                                                        DNA;
                                                                             exon; gene expression analysis;
                                                                                                    exon
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WO200157275-A2

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AAK33709/C
ID AAK327
XX
AC AAK327
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DT 06-NOV
DT 06-NOV
DE Human
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27-SEP-2000;
04-OCT-2000;
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26 - MAY - 2000;
30 - JUN - 2000;
03 - AUG - 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is one of the probes of the
                                                                                                                  Human; bone
microarray;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The
                                            WO200157276-A2
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                                                                                                                                                                                                                                                  AAK32709;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 475 BP;
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                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                               GAGGAGGAG 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CAGAGCCAGCTCCCTGAGGTGGAGGAGAGTGATCCGAGACATGGGTGTGGGACAGTCAGCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAGGAGGCGGGCCGGCTCAGGAGCAAGATGAAGACCATGGAGCAGCAGATTGAGCTTCTACTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100;
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                                                                                                                                                                          bone marrow
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                                                                                                                                                                                                                                                                                      standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ
                                                                                                                  cancer;
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ilarity 52.9%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0234687.
2000US-0236359.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic
                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001WO-US00667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           38
                                                                                                                                                                        expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NO:
                                                                                                                                   expressed
                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ą
                                                                                                                    leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        provides a number of single exon nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6964;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         198 C;
                                                                                                                                                                                                                                                                                      475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Σ
                                                                                                                                                                                                                                                                                      ЧB
                                                                                                                                                                      single exon probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           650pp +
                                                                                                                exon; gene expression
a; lymphoma; myeloma; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 46.6;
Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>ن</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           194 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence Listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 analyzing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .018;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene
                                                                                                                                                                      SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                        ID NO:
                                                                                                                                   analysis; probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English
                                                                                                                                                                        7266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             245
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RESULT 20
AAI38524/c
ID AAI385
XX AAI385
AC AAI385
AC AAI385
AC Probe
DF 17-OCT
XX
Probe
CX
DF Probe
XX
Probe;
KW Probe;
KW Geneti
XX
GS Homo s
XX
PN WO2001
AX
PF 30-JAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Qy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Вb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Qy
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                                                                                                                                                                                                                                                                                                                                                                                                                184
                                                                                                                                                                                                                                                                                                                                                                                                                                                     682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      562
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Query Match
Best Local Sin
Matches 100;
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26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancer such as lymphoma, leukaemia and myeloma. The present sequence is one of the invention.
30-JAN-2001; 2001WO-US00663.
                         09-AUG-2001
                                                 WO200157272-A2
                                                                                                   genetic
                                                                                                                                       Probe #7210 used to measure gene expression in human placenta
                                                                                                                                                                 17-OCT-2001
                                                                                                                                                                                          AAI38524;
                                                                                                                                                                                                                  AAI38524 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human genome-derived single exon nucleic acid analyzing gene expression in human bone marrov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides a number of single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                     GAGGAGGCGGGCCCGGCTCAGGAGCAAGATGAAGACCATGGAGCAGATTGAGCTTCTACTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACACTGAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAA 561
                                                                           sapiens
                                                                                                                                                                                                                                                                                   GAGGAGGAG
                                                                                                                                                                                                                                                                                                           GTGGAACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                        AAACAGAAGAAGAAGAAGAAGGAAGGAGGAGGAGGAGGAAGGAAGAAGAAA 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001-488900/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SG,
                                                                                                                                                                                                                                                                                                                                   CAGAGCCAGCTCCCTGAGGTGGAGGAGATGATCCGAGACATGGGTGTGGGACAGTCAGCG
                                                                                                                                                                                                                                                                                                                                                                                      microarray;
c disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULAR DYNAMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP;
                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001WO-US00668
                                                                                                                                                                                                                                                                                                            690
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                                                                                                     SS
                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.3%;
                                                                                                              human; placenta; antenatal diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7266; 658pp + Sequence Listing;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Score 46.6; D; Pred. No. 0.01: 0; Mismatches
                                                                                                                                                                                                                    ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       46.6;
No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   marrow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .018;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             probes useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                       sample
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one of
                                                                                                                                                                                                                                                                                                                                    185
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RESULT 21
ABA71159/c
ID ABA71159;
XX
AC ABA71159;
XX
DE Human foetal liver s
XX
DE Human; foetal liver;
XX
DE Human; foetal liver;
XX
DE Homo sapiens.
XX
PN WO200157277-A2.
XX
PD 09-AUG-2001.
XX
PF 30-JAN-2000; 2000US-PR 26-MAY-2000; 2000US-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 100
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26-MAY-2000; 2000US-0208408.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0234539.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to single exon nucleic acid probes The present sequence is one suppose. The probes are useful for producing a microarray for predicting, measuring and displaying expression in samples derived from human placenta. The probes are for antenatal diagnosis of human genetic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 475 BP; 38 A; 198 C; 45 G; 194 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Penn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   622
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          562
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Sir
hes 100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAGGAGGAG 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTGGAACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAGAGCCAGCTCCCTGAGGTGGAGGAGATGATCCGAGACATGGGTGTGGGACAGTCAGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAGGAGGCGGGCCGGCTCAGGAGCAAGATGAAGACCATGGAGCAGATTGAGCTTCTACTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001-488897/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genome-derived single exon nucleic aci
zing gene expression in human placenta
                                                                                                                                                                                                                              foetal liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULAR DYNAMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hanzel DK,
     ; 2000US-0180312.
; 2000US-0207456.
; 2000US-0608408.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                              2001WO-US00669
                                                                                                                                                                                                                                                                  liver single
                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          690
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                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                     entry)
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                                                                                                                                                                                                                              gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chen
                                                                                                                                                                                                                                                                                                                                                                               511
                                                                                                                                                                                                                              expression;
                                                                                                                                                                                                                                                                  exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           654pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Σ
                                                                                                                                                                                                                                                                                                                                                                               ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 46.6; DB Pred. No. 0.018; 0; Mismatches
                                                                                                                                                                                                                                                                  nucleic acid probe #19464.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
an placenta -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
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                                                                                                                                                                                                                              single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
                                                                                                                                                                                                                              exon
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                                                                                                                                                                                                                              nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 475;
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probes are useful
                                                                                                                                                                                                                              acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                            probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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 RRESULT Z
ABA37497,
ID ABA3
XX ABA3
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XX Prob
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XX Huma
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KW Card
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XX Hom
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Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format direction wipo at ftp.wipo.int/pub/published_pct_sequences.
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21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                      congenital heart
                                                                                                                                                                                                                                                                                                        Human; gene expression; cardiovascular disease;
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30-JAN-2001; 2001WO-US00666
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                                                                                                                 microarray; Alzheimer's disease;
epilepsy; cancer; ss.
                                                                                                                                 Human; brain expressed exon; gene expression analysis; probe; microarray; Alzheimer's disease; multiple sclerosis; schizophrenia;
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RESULT 24
AAK45444/C
ID AAK454
XX
AC AAK454
XX
DT 06-NOV
DT 06-NOV
XX
DE Human
XX
KW Human;
KW microa
XX
CS Homo s
XX
PN WO2001
XX
PD 09-AUG
                                                                                                                                                                                                       밁
                                                                                                                                                                                                                       QY
                                                                                                                                                                                                                                              Вb
                                                                                                                                                                                                                                                                Qy
                                                                                                                                                                                                                                                                                      В
                                                                                                                                                                                                                                                                                                         Š
                                                                                                                                                                                                                                                                                                                             В
                                                                                                                                                                                                                                                                                                                                               δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.3%;
Best Local Similarity 52.9%;
Matches 100; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is one of the probes of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Single
brains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Penn
                                                           microarray;
                                                                       Human;
                                                                                                             06-NOV-2001
                                                                                                                                                    AAK45444 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-AUG-2001
                                                                                                                                                                                                                                                                                                                            267
                                                                                                                                                                                                                                                                                                                                                                                                                                            invention.
                                                                                                                                                                                                                                               147
                                                                                                                                                                                                                                                                                                                                         502 ACACTGAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAA 561
                                                                                                                                                                                                                         682 GTGGAACAG
                                                                                                                                                                                                                                                                                       207
                                                                                                                                                                                                       87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention provides a number of single exon nucleic acid
                                                                                                                                                                                                                                                                                                                             GAGGAGGAG
                                                                                                                                                                                                                                               CAGAGCCAGCTCCCTGAGGTGGAGGAGATGATCCGAGACATGGGTGTGGGACAGTCAGCG
                                                                                                                                                                                                                                                                                      GAGGAGGCGGCCGGCTCAGGAGCAAGATGAAGACCATGGAGCAGATTGAGCTTCTACTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001-483446/52.
                                                                                        bone marrow expressed single exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      exon
                                                                     bone marrow expressed exon; gene expression analysis; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULAR DYNAMICS
                                                                                                                                                                                                                                                                                                                                                                                                                        511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hanzel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
2000US-0024263.
                                                            cancer;
                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0180312
2000US-0207456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001WO-US00667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IJ
                                                                                                                                                                                                       79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NO:
                                                                                                                                                                                                                                                                                                                                                                                                                        Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acid
                                                            leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19446;
                                                                                                                                                                                                                                                                                                                                                                                                                        231 C;
                                                                                                                                                      511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ٤
                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                              Score 46.6; DB Pred. No. 0.019;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          650pp +
                                                                                                                                                                                                                                                                                                                                                                                                                        26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rank
                                                            Lymphoma;
                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for
                                                                                                                                                                                                                                                                                                                                                                                                                        <u>و</u>;
                                                                                                                                                                                                                                                                                                                                                                                                                        235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      analyzing gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                        T; 0
                                                                                         probe SEQ ID NO:
                                                            myeloma;
                                                                                                                                                                                                                                                                                                                                                                                        22;
                                                                                                                                                                                                                                                                                                                                                                      ; 68
                                                                                                                                                                                                                                                                                                                                                                                                                        other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Listing;
                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English
                                                                                                                                                                                                                                                                                                                                                                                          511;
                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   samples
                                                                                                                                                                                                                                                                                     148
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WO200157276-A2

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RESULT 25
AA151389/C
ID AA1513
XX
AA151389/C
ID AA1513
XX
Probe
DE Probe
XX
Prob

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04 FEB 2000;
26 MAY 2000;
30 JUN 2000;
03 AUG 2000;
21 SEP 2000;
27 SEP 2000;
04 OCT 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukaemia and myeloma. The present sequence is one of the invention.
                                                                                                                                                                                                                                                                                            17-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Penn
    30-JAN-2001; 2001WO-US00663
                                                  09-AUG-2001
                                                                                                                                                                                genetic
                                                                                                                                                                                                     Probe; microarray; human;
                                                                                                                                                                                                                                                   Probe #20075
                                                                                                                                                                                                                                                                                                                                          AAI51389;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JAN-2001;
                                                                                           WO200157272-A2
                                                                                                                                                                                                                                                                                                                                                                                   AAI51389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     502 ACACTGAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GTGGAACAG 690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAGAGCCAGCTCCCTGAGGTGGAGGAGATGATCCGAGACATGGGTGTGGGACAGTCAGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAGGAGGCGGCCGGCTCAGGAGCAAGATGAAGACCATGGAGCAGATTGAGCTTCTACTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAGGAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001-488900/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genome-derived single exon nucleic acid probes useful
zing gene expression in human bone marrow -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4
                                                                                                                                                                                   disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632368.
2000US-0234687.
2000US-02346359.
2000US-0236359.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention
                                                                                                                                                                                                                                                 used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 19 A;
                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001WO-US00668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               79
                                                                                                                                                                                   SS
                                                                                                                                                                                                                                              to measure gene expression in human placenta sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  provides a number of single exon nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20001; 658pp + Sequence Listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               231 C;
                                                                                                                                                                                                                                                                                                                                                                                   511
                                                                                                                                                                                                     placenta; antenatal diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ξ
                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 46.6; DB Pred. No. 0.019;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            511;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            208
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  621
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
RESULT 2
AADOGTO
XX AADOG
XX AADO
XX AADO
DT 06-8
DT 06-8
XX Huma
XX Huma
XX Huma
XX Huma
XX Homa
XX ACO
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ър
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Вb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DЬ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ъ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Penn
                                                                                                                                                                                                                          acute
                                                                                                                                                                                                                                                                                                               Human
                                                                                                                                                                                                                                                                                                                                                          06-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                      AAD06778;
                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD06778 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  analyzing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             87 GAGGAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GTGGAACAG
                                                                                                                                                                                sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001-488897/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SG
                                                                                                                                                                                                                       myeloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULAR DYNAMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hanzel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
2000US-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                       leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA;
                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chen
                                                                                                                                                                                                                       drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                 4181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ξ
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The present invention relates to single exon nucleic acid probes the present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying get expression in samples derived from human placenta. The probes are for antenatal diagnosis of human genetic disorders.
                                                                                                                                                                     Human; haematopoietic lineage switch; HLS-5; tumour suppressor;
ring finger B-box coiled-coil; RBCC; transcriptional regulator;
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zing gene expression in human placenta -
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/product= "Human HLS-5
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                             CDS
                                                                                                Human; haematopoietic lineage switch; HLS-5; tumour suppressor; ring finger B-box coiled-coil; RBCC; transcriptional regulator; acute myeloid leukaemia; drug screening; gene therapy; cytostati
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                                                     nootropic; neuroprotective; anticonvulsant; antiparkinsonian; anti-HIV; antidiabetic; immunostimulant; immunomodulator; antilinflammatory; antithyrodid; immunosuppressive; nephrotropic; antigout; thyromimetic; cytostatic; antibacterial; virucide; fungicide; protozoacide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a tumour suppressor gene, termed haematopoietic lineage switch (HLS)-5 and the protein it encodes. HLS-5 is a member of ring finger B-box coiled-coil (RBCC) family of
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                                                                                                                                         Human RECAP polynucleotide,
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                                         antiarteriosclerotic;
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07-OCT-1999;
12-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease, dementia, Parkinson's disease, Downs's syndrome, amyotrophic lateral sclerosis, multiple sclerosis, bacterial and viral meningitis, CJD (Creutzfeldt-Jakob disease), GSS (Gerstmann -Straussler-Scheinker syndrome); immunological disorders, including autoimmune/inflammatory disorders such as AIDS, DiGeorge's syndrome, severe combined immunodefictency disease (SCID), Chediak-Higashi syndrome, Cushing's disease, Addison's disease, autoimmune thyroiditis, Crohn's disease, diabetes mellitus, Good pasture's syndrome, gout, Grave's diseases, Hashimoto's thyroiditis, Sjogren's syndrome, Werner's syndrome, viral, bacterial, fungal, parasitic, protozoal, and helminthic infections; an cell proliferation disorders such as arteriosclerosis, atherosclerosis
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Baughn MR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel receptors and associated proteins for diagnosis and treatment of neurological disorders, immunological disorders including autoimmune/inflammatory disorders and cell proliferative disorders such as cancer
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n C, Lal P;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       other;
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507 GAAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAAGAGGA 566

Best Local Similarity Matches 132; Conserv

Conservative

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Score 45; DB 2 Pred. No. 0.095 0; Mismatches

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Length 1824;

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Query Match Best Local

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RESULT 29
AAS81488/c
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                         for identifying expressed genes. (I) is useful in gene therapy techniques (II) (II) (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations cresponsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. AAS64197-AAS94564 represent novel human diagnostic coding sequences of the invention.

Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at first wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                        The invention relates to isolated polynucleotide (I) and polyperide (II) sequences. (I) is useful as hybridisation probes, polyperase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifiers.
                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polynucleotide and encoded polypeptides, us diagnostics, forensics, gene mapping, identification of responsible for genetic disorders or other traits and to biodiversity
 Sequence 1824 BP;
                                                                                                                                                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-639362/73.
P-PSDB; ABG17301.
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23-AUG-2000; 2000US-0649167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA encoding novel human diagnostic protein #17292.
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                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID No 17292; 103pp; English
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270 A; 670 C;
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RESULT 3
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                                       The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity.
                                                                                                                                                                                                                                                                                       New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity -
                The polypeptide and pol diagnostics, forensics,
                                                                                                                                                                                                                                                        Claim
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23-AUG-2000;
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DB; ABG15508.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             HYSEQ INC
tide and polynucleotide sequences have applications, forensics, gene mapping, identification of mutatic for genetic disorders or other traits to assess bio
                                                                                                                                                                                                                                                                                                                                                                                                              Liu
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2000US-0649167.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 e mapping; gene mapping; gene therapy; forensic;
medical imaging; diagnostic; genetic disorder;
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RESULT 31
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Best Local S
Matches 132
                                                                                                                                                                                                                                                    04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
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Note: The sequence data for this patent did not appear in the prispecification, but was obtained in electronic format directly from
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                                                                                                                                                                           (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001; 2001WO-US00667
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human brain expressed single exon probe
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                                                                         2001-483446/52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer;
                                                                                                                           Hanzel
                                                                                                                                                                                                                          ; 2000US-0207456.
; 2000US-0608408.
; 2000US-0632366.
; 2000US-0234687.
; 2000US-02346359.
; 2000US-0236359.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                   2000US-0180312
                                                                                                                         DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            267
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
Pred.
                                                                                                                           Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             937 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           45;
No.
                                                                                                                           DR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               474 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NO: 19590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 2850;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
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from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and
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Single exon nucleic acid brains -

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AAK45604/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
  WPI;
                                          Penn
                                                                                                                                                                                                                                      04-FEB-2000;
26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia,
                                                                                                                                                                                            30-JUN-2000;
03-AUG-2000;
                                                                                                                                                                                                                                                                                                       30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                  09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                          WO200157276-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           microarray; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; bone marrow expressed exon; gene expression analysis; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human bone marrow expressed single exon probe SEQ ID NO: 20161.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK45604;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            epilepsy and cancers. The present sequence is one of the probes of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAK45604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probes which are derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4;
                                                                                     (MOLE-)
                                                                                                                                                   27-SEP-2000;
                                                                                                                                                                          21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           588
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      648
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Si
hes 131;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19
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2001-488900/53
                                          SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAAGAAGAAAAGAAGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTCTCTCAAGAAAGAGTACGAGAATCTAAAAGAGGCACGGAAGGCCTCAGGGGAGGTGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GATGAAGACCATGGAGCAGATTGAGCTTCTACTCCAGAGCCAGCTCCCTGAGGTGGAGGA 647
                                                                                     MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
                                          Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ
                                                                                                                           ; 2000US-0180312.
; 2000US-0207456.
; 2000US-0608408.
; 2000US-0632366.
; 2000US-0234687.
; 2000US-0234687.
; 2000US-0234683.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                       2001WO-US00668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 151 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.2%;
                                          Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19590; 650pp + Sequence Listing; English
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Pred. No. 0.054;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lymphoma; myeloma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 109
                                          Rank
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      126;
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Best Local Similarity
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                                                                                          31-MAR-2000;
23-AUG-2000;
                   P-PSDB; ABG10053
                                                                     (HYSE-) HYSEQ
                                                                                                                         30-MAR-2001; 2001WO-US08631
                                                                                                                                              11-OCT-2001
                                                                                                                                                                  WO200175067-A2
                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                    Human; chromosome mapping; gene mapping; gene therapy;
                                                                                                                                                                                                                                       DNA encoding novel
                                                                                                                                                                                                                                                              13-FEB-2002
                                                                                                                                                                                                                                                                                                      AAS74240 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; SEQ ID NO: 20161; 658pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      analyzing gene expression in human bone marrow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human genome-derived single exon nucleic acid probes useful
                                                                                                                                                                                                                                                                                                                                                                                                                     708
                                                                                                                                                                                                                                                                                                                                                                                                                                          139
                                                                                                                                                                                                                                                                                                                                                                                                                                                               648
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                                                                                                                                                                                                                                                                                                                                                                                                  79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention
                             2001-639362/73
                                                                                                                                                                                                          supplement; medical imaging; diagnostic; genetic
                                                                                                                                                                                                                                                                                                                                                                                                              GTCTCTAAGAAAGAGTACGAGAATCTAAAAGAGGCCCGGAAGGCCTCAGGGGAAGGTGGC 767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGAGGAAAAGGAGAAGGAGGAGAAACAGGTGGAGGAGGAGGAGGAGGAGGAGGAGGAG 199
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                                                                                                                                                                                                                                                                                                                                                                                                131;
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                                                                                                                                                                                                                                                             (first entry)
                                                                                          2000US-0540217.
2000US-0649167.
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                                                 Ç
                                                                                                                                                                                                                                        human diagnostic protein #10044.
                                                                                                                                                                                                                                                                                                      cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 44.4;
Pred. No. 0
                                                                                                                                                                                                                                                                                                       ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     109 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .054;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treatment of cancers sequence is one of
                                                                                                                                                                                                          disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
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                                                                                                                                                                                                           SS
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New isolated polynucleotide and encoded polypeptides, useful in

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RESULT 3
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DI 13-F
DI DNA
XX Huma
KW Huma
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OS Homc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC The invention relates to isolated polynucleotide (I) and CC polypeptide (II) sequences. (I) is useful as hybridisation probes, CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome CC and gene mapping, and in recombinant production of (II). The CC polynucleotides are also used in diagnostics as expressed sequence tags CC for identifying expressed genes. (I) is useful in gene therapy techniques CC to restore normal activity of (II) or to treat disease states involving CC (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical CC imaging of sites expressing (II). (I) and (II) are useful for treating CC disorders involving aberrant protein expression or biological activity. CC The polypeptide and polynucleotide sequences have applications in CC diagnostics, forensics, gene mapping, identification of mutations cresponsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and CC amino acid sequences. AAS64197-AAS94564 represent novel human CC diagnostic coding sequences of the invention.

CC Note: The sequence data for this patent did not appear in the printed coast first pube introuble and create and products format directly from WIPO as the polynomial sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 121; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID No 10044; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnostics,
                  WO200175067-A2
                                                                                                                                         DNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 693 BP; 296 A; 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 responsible for
                                                   Homo sapiens
                                                                                                        Human;
                                                                                                                                                                            13-FEB-2002
                                                                                                                                                                                                               AAS90715;
                                                                                                                                                                                                                                                  AAS90715 standard; cDNA; 693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          502 ACACTGAAAAAGCAGATGAAGTACTTAGAGCAGCAGGATGAGATGAGACCAAACAAGCACAA 561
                                                                                                                                                                                                                                                                                                                                        448
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                                                                                                                                                                                                                                                                                                                                                                                                                               GTGGAACAGCTGGCTGTGTGTGTGTCTCTCAAGAAAGAGTACGAGAATCTAAAAGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CAGAGCCAGCTCCCTGAGGTGGAGGAGATGATCCGAGACATGGGTGTGGGACAGTCAGCG
                                                                                                                                                                                                                                                                                                                                      gagaagaag
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                                                                                                                                                                                                                                                                                                                                                                                                         chromosome
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                                                                                                                                      novel human diagnostic protein #26519
                                                                                                                                                                          (first entry)
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                                                                                    medical imaging; diagnostic;
                                                                                                     mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.2%;
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                                                                                                      gene mapping;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               234 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                     gene therapy;
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                                                                                    genetic
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                                                                                      disorder;
                                                                                                      forensic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-MAR-2000;
23-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ocal Similarity
                                                                                                                                                                                                                                                                                                   GAGGAGGCGGGCCGGCTCAGGAGCAAGATGAAGACCATGGAGCAGATTGAGCTTCTACTC
                                                                                                                 CAGAGCCAGCTCCCTGAGGTGGAGGAGATGATCCGAGACATGGGTGTGGGACAGTCAGCG
                                                                                                                                                                                                                                                                                                                                                                   2001-639362/73
gagaagaag
                                               GCACGGAAG 750
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP;
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2000US-0649167.
  456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 44.2; DB Pred. No. 0.097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          234 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               128;
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Best Local Similarity
Matches 127; Conserv
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29-DEC-1995;
29-DEC-1995;
30-JAN-1996;
30-JAN-1996;
                                                                                                                                                                                                                                                        This sequence represents a fragment of the genomic sequence containing the coding region for the mouse WRN gene (AAR38004). The corresponding human gene (AAR3001) encodes a protein related to Merner's syndrome. The products can be used for the detection and treatment of Merner's syndrome (WS), an autosomal recessive disorder with a complex phenotype.
                                                                                                                                                                                                                                                                                                                                          Isolated nucleic acid molecule encoding the WRN gene product useful for detection and treatment of Werner's syndrome, and
                                                                                                                                                                                                                                 Sequence 51259 BP; 14533 A;
                                                                                                                                                                                                                                                                                                                Claim 1; Fig 7; 153pp; English.
                                                                                                                                                                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                                                                                                                                                          Fu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Partial mouse WRN genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-JUL-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     recessive
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                                                       989
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                                                                                                                                                                                                                                                                                                                                                                                                             (/IHSO)
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                    746
                                                                                                             207
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                                     87
   27
                                                                                                                                                                                                                                                   well as related diseases
            GGAAGGCCTCAGGGGAGGTGGCTGA
                                                                                  GCCAGCTCCCTGAGGTGGAGGAGAGATGATCCGAGACATGGGTGTGTGGGACAGTCAGCGGTGG
                                                                                                                                                                  TGAAAAAGCAGATGAAGTACTTAGAGCAGCAGCAGGATGAGACCAAACAAGCACAAGAGG
                                                                                                                             AGGCGGGCCGGCTCAGGAGCAAGATGAAGACCATGGAGCAGATTGAGCTTCTACTCCAGA 625
                                                                                                                                                AACAGCTGGCTGTGTACTGTGTGTCTCTCAAGAAAGAGTACGAGAATCTAAAAGAGGCAC
                                                                         WRN; Werner'
ive disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                            OSHIMA .
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                                                                                                                                                                                    Conservative
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95US-0009409.
95US-0580539.
96US-0010835.
96US-0594242.
                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULAR CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96WO-US20785
                                                                                                                                                                                           2.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA; 51259
                                                                                                                                                                                                                                                                                                                                                                                          Oshima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    phenotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             syndrome;
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                                                                                                                                                                                                                                 9635 C;
                                                                                                                                                                                    0;
                                                                                                                                                                                           Score 44.2;
Pred. No. 0.
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                                                                                                                                                                                                                                                                                                                                                                                         Schellenberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     detection; ss.
                                                                                                                                                                                                                                10266 G;
                                                                                                                                                                                            DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis; autosomal;
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Best Local Similarity
Matches 128; Conserv
                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invention useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA sequences (ABL101840-ABL16175) and the encoded proteins (ABB5737-ABB72072).
                        1873
                                                                      1813
                                                                                                                     1753
                                                                                                                                                                   1693
                                                                                                                                                                                                                                                                                               The sequence data for this patent did not form specification, but was obtained in electronic at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic genes from Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-656860/75
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11-JUL-2000; 2000US-0614150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interactions
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                                                                                                                                463 CTGCAGCAGGCCTTGGGCAAGGCCGAGATGCTGTGCTCCACACTGAAAAAAGCAGATGAAG 522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PEKE
                                                                                                                                                                                         403 CAGGTCATCATCGACACTCTGCGGGATACGCTGGAAGAACGCAATGCTACTGTGGTATCT 462
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GAGGAGATGATCCGAGACATGGGTGTGG
                                                                                                                                                                 cagcacaccaaggctgaactgctggacaagcggaaggagaacgatcggcacgccaagtcg
                      cacagccaggagaacctgcaaacggtgtacgatcagacccagaggcagtgcagccagtat
                                            AGCAAGATGAAGACCATGGAGCAGATTGAGCTTCTACTCCAGAGCCAGCTCCCTGAGGTG
                                                                    ctgcaagaggacgagcagaagctggcgcgcctgcgcaaggatgtggacaacctgatgaac
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                                                                                                                                                                                                               Score 44; DB:
Pred. No. 0.21
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                                                                                                                                                                                                                                                                                                                                                         showed the inv(16) fusion to comprise a sequence from the CBFB gene, encoding a novel transcription factor, and the MYHI1 gene, encoding smooth muscle myosin heavy chain. In 1 patient, nt 1-492 of the CBFB gene were fused to nt 994 of MYHI1 (shown in AAQ84589; predicted as sequence in AAR66930). Probes specific for inv(16) c
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR was performed on total cellular RNA from 5 AMML patients having a pericentric inversion of chromosome-16, M4Eo subtype. Sequencing showed the inv(16) fusion to comprise a sequence from the CBFB
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DB; AAR66930.
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gccaggccaagcaggaggtggaacataagaagaagaagctggaggcgcaggtgcaggagc
                                          TGAAGACCATGGAGCAGATTGAGCTTCTACTCCAGAGCCAGCTCCCTGAGGTGGAGGAGA 649
                                                                                      agaataagcagacgctggagaaagagaacgcagacctggccggggagctgcgggtcctgg
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6); CBF-beta; CBFB gene; transcription factor; myosin; MYH11;
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UNIV TEXAS SYSTEM.
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                                                             The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The inventiuseful in developmental biology and in elucidating cell signalling cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABLJ01840-ABLJ6175) and the encoded proteins sequences (ABLJ01840-ABLJ6175) and the encoded proteins
                                      The sequence data for this patent did not form specification, but was obtained in electronic f
                                                                                                                                                                                  Claim 1; SEQ ID NO 24932; 21pp + Sequence Listing; English
                                                                                                                                                                                                             genes from Drosophila and interactions -
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11-JUL-2000; 2000US-0614150
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                           at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                      New isolated nucleic acid
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                                                                                                          The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invent useful in developmental biology and in elucidating cell signalling cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABLIG176-ABL30511), expressed DNA sequences (ABLG1840-ABL16175) and the encoded proteins
                                                                          The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                               New isolated nucleic acid genes from Drosophila and
                                                                                                                                                                                                                 Claim 1;
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11-JUL-2000;
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pharmaceutical; gene; ss.
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Query Match Best Local Similarity

2.2%;

Score Pred.

No.

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Length 16442;

Sequence as well as

16442 BP;

4392 A; 2975 diseases

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RESULT 40
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ID AAX830
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29-DEC-1995;
29-DEC-1995;
30-JAN-1996;
30-JAN-1996;
                                                                      This sequence represents a fragment of the genomic sequence containing the coding region for the mouse WRN gene (AAR83004). The corresponding human gene (AAR3001) encodes a protein related to Werner's syndrome. The products can be used for the detection and treatment of Werner's syndrome (WS), an autosomal recessive disorder with a complex phenotype
                                                                                                                                                                                                                    Claim 1; Fig 7;
                                                                                                                                                                                                                                                                           diseases
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Search completed: September 4, 2002, 14:00:32 Job time: 9036 sec
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                                                                                                                                                                           632 TCCCTGAGGTGGAGATGATCCGAGACATGGGTGTGGGACAGTCAGCGGTGGAACAGC 691
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                                                                                                                                                                                                                                                                                                               0; Mismatches 144; Indels
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